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TECH CENTER 1600/2300

SEQUENCE LISTING

(1) GENERAL INFORMATION

- (i) APPLICANT: ISNER, Jeffrey
ASAHARA, Takayuki
- (ii) TITLE OF THE INVENTION: METHODS FOR REGULATING ANGIOGENESIS
(as amended)
- (iii) NUMBER OF SEQUENCES: 2
- (iv) CORRESPONDENCE ADDRESS:
(A) ADDRESSEE: DIKE, BRONSTEIN, ROBERTS & CUSMAN, LLP
(B) STREET: 130 Water Street
(C) CITY: Boston
(D) STATE: MA
(E) COUNTRY: USA
(F) ZIP: 02109
- (v) COMPUTER READABLE FORM:
(A) MEDIUM TYPE: Diskette
(B) COMPUTER: IBM Compatible
(C) OPERATING SYSTEM: DOS
(D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION DATA:
(A) APPLICATION NUMBER: 08/744,882
(B) FILING DATE: 08-NOV-1996
(C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
(A) APPLICATION NUMBER:
(B) FILING DATE:
- (viii) ATTORNEY/AGENT INFORMATION:
(A) NAME: Resnick, David S
(B) REGISTRATION NUMBER: 34,235
(C) REFERENCE/DOCKET NUMBER: 46963
- (ix) TELECOMMUNICATION INFORMATION:
(A) TELEPHONE: 617-523-3400
(B) TELEFAX: 617-523-6440
(C) TELEX:

(2) INFORMATION FOR SEQ ID NO:1:

- (i) SEQUENCE CHARACTERISTICS:
(A) LENGTH: 30 base pairs
(B) TYPE: nucleic acid
(C) STRANDEDNESS: single
(D) TOPOLOGY: linear

49

B

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AAGACATTTT CGGGCTCAG CTGCGCACCC

30

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TGGGGTAGGC ACTTTAGTAG TTCTCCTAAC

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Cont

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